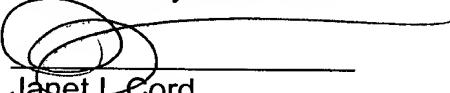


In the Specification

Page 6, paragraph 1, please replace as follows:

Figure 5 shows an amino acid sequence alignment for the PHD finger motif of AIR-1 (SEQ ID NO:3 and SEQ ID NO:4), Mi-2 (SEQ ID NO:5 and SEQ ID NO:6), and TIF1 (SEQ ID NO:7). The consensus amino acid residues conserved in the PHD finger motif is indicated by the bold letters underneath the residues that are identical with AIR-1 (aa 299-340) (SEQ ID NO:3) are shown by the dots. GenBank accession nos. of Mi-2 and TIF1 are X86691 and AF009353, respectively.

Respectfully submitted


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